



NCBI Resources for NextGen Sequence Analysis

Aligning NextGen data with magic-blast and examining the result in Genome Data Viewer

<https://go.usa.gov/xVtK7> | <https://www.ncbi.nlm.nih.gov/genome/gdv/>

National Center for Biotechnology Information • National Library of Medicine • National Institutes of Health • Department of Health and Human Services

Magic-BLAST

A standalone commandline variant of NCBI's famous BLAST sequence search algorithm, this tool was designed for alignment of DNAseq and RNA seq data to a genomic assembly and the results can be stored in several formats.

- **Magic-BLAST Homepage:** <http://bit.ly/2M6BI01>
- **Magic-BLAST Publication:** <https://go.usa.gov/xVtsM>

Genome Data Viewer

The NCBI Genome Data Viewer (GDV) is an interactive genome browser for the exploration and analysis of many types of sequence data in context with curated annotations on an eukaryotic RefSeq genome assembly. This tool facilitates the examination of data from NCBI resources, track hubs, or your own data sets which can be incorporated by uploading a file or streaming your data from a remote location with a URL.

- **GDV Homepage:** <https://go.usa.gov/xVtsj>
- **GDV Documentation:** <https://go.usa.gov/xVtsD>



Human Reference Genome Sequences

Magic-BLAST aligns nextgen data to a reference genomic assembly. The Human Reference Genomic Assembly produced by the Genome Reference Consortium (GRC) is a standard for the genomic research community and this assembly is annotated by the NCBI Reference Sequences (RefSeq) project to facilitate understanding and comparison of the data.

- **GRC Homepage for Human:** <https://go.usa.gov/xVts9>
- **NCBI RefSeq Homepage:** <https://go.usa.gov/xVtsn>
- **Human Genome Resources Page:** <https://go.usa.gov/xVtsP>

NextGen Sequence Reads

A source of nextgen data for analysis with Magic-BLAST and visualization with GDV is the NCBI Sequence Read Archive (SRA) database. SRA sequence reads can be downloaded from the NCBI website and are also now available in the AWS and GCP clouds.

- **SRA Homepage:** <https://go.usa.gov/xVtHg>
- **SRA Search & Download Documentation:** <https://go.usa.gov/xVtH4>
- **SRA in the Cloud Documentation:** <https://go.usa.gov/xVtHD>

An Example of Using NCBI's NextGen Sequence Analysis Tools

Finding NextGen Sequence Data at NCBI

The SRA database provides access to submitted nextgen sequence datasets. These can be found by searching SRA directly or from a PubMed record which describes the dataset in a publication.

- **PubMed Record:** <https://go.usa.gov/xVtAd>

The PubMed SRA link in the “Related Information” section provides a quick way of finding the relevant SRA data records. To download the SRA sequence data, you can use either the SRA Run browser, SRAToolKit or access the data in the AWS or GCP Clouds.

- **SRA Record:** <https://go.usa.gov/xVtAs>
- **SRA Download Tips:** <https://go.usa.gov/xVtA6>

Aligning NextGen Sequence Data with Magic-BLAST

The Magic-BLAST GitHub site has guidance on how to download to your own computer, preparation of a reference assembly for alignment, and use of the application with SRA and your own data. Please note that to visualize Magic-BLAST alignments in GDV, you will need to convert the SAM to BAM format which can be done with several open source tools, including SAMtools.

- **Download Magic-BLAST:** <http://bit.ly/2IHQn69>
- **Making a BLAST Database:** <http://bit.ly/2M5dxFQ>
- **SRA as Input to Magic-BLAST:** <http://bit.ly/329ZkNp>

Examining NextGen Sequence data with the Genome Data Viewer

The GDV tool provides both written tutorials and a YouTube video tutorial playlist to assist users in learning to utilize the features and functionality of this versatile visualization tool.

- **GDV Written Tutorials:** <https://go.usa.gov/xVtHs>
- **GDV YouTube Playlist:** <http://bit.ly/35tikIE>

GDV provides portlets to “Search” the genomic assembly to zoom-in to a region of interest. “The User Data and Track Hubs” provides a simple way to “Add Remote Files” such as Magic-BLAST alignments. Then, full functionality of GDV enables comparison with curated and annotated data by configuring the data “Tracks” shown.

Example Magic-BLAST alignments:

- <https://go.usa.gov/xVtHz>
- <https://go.usa.gov/xVtHh>

The collage illustrates the workflow from finding SRA data in a PubMed record to visualizing it in the Genome Data Viewer. It shows the SRA Run browser interface, the SRA archive data table, and the GDV interface with various tracks and a detailed view of a specific genomic region.

Type	Size	Location	Name	Free Egress	Access Type
run	4,787,814 Kb	NCBI	https://sra-download.ncbi.nlm.nih.gov/traces/sra47/SRR005325/SRR5452960	worldwide	anonymous
		AWS	s3://sra-pub-run-2/SRR5452960/SRR5452960.1	s3.us-east-1	aws identity
		GCP	gs://sra-pub-run-2/SRR5452960/SRR5452960.1	gs.us	gcp identity